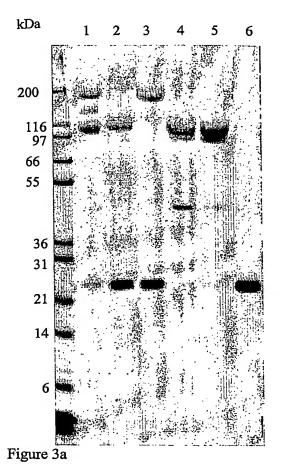


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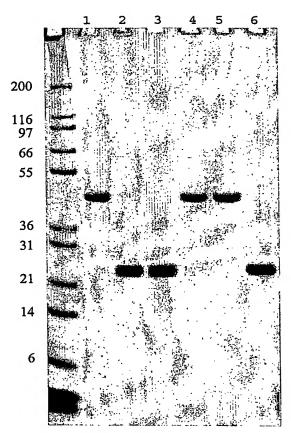
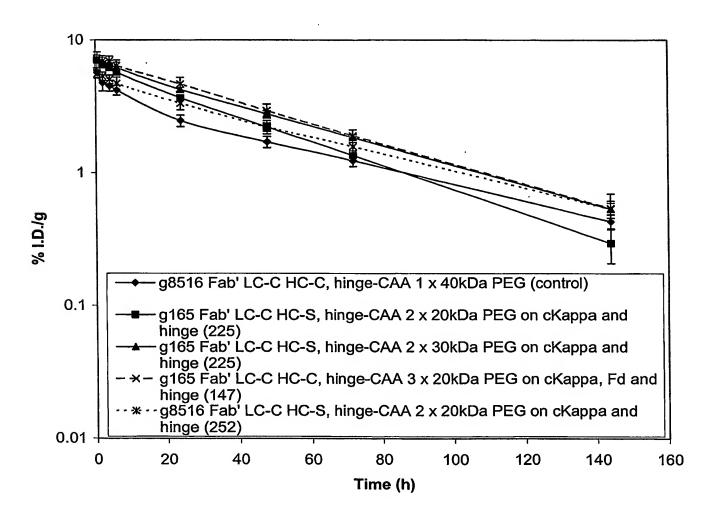


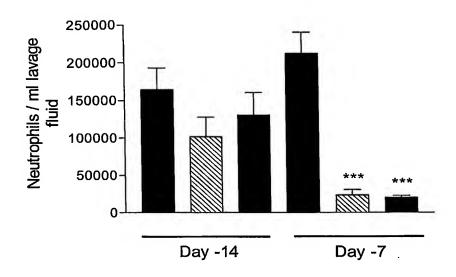
Figure 3b

Figure 4



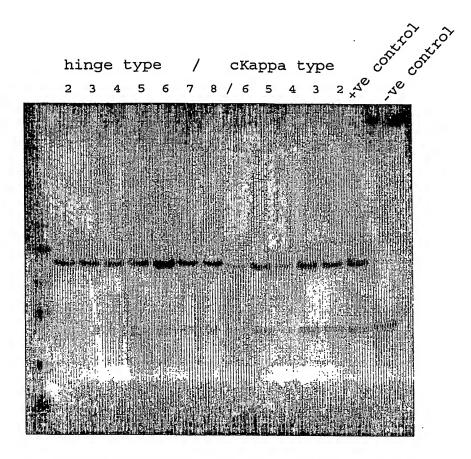
WO 2005/003171 PCT/GB2004/002871 5/8

Figure 5



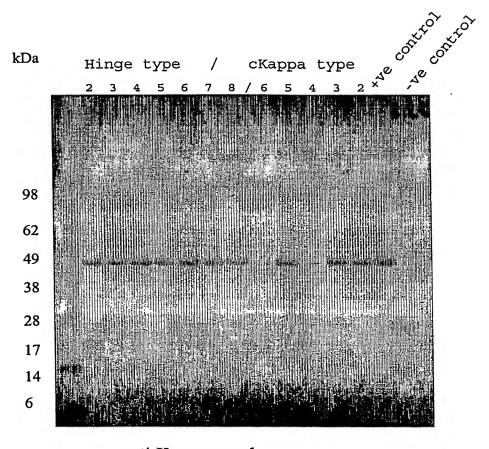
gA33(LC-C,HC-C, hinge-CAA)Fab'-PEG(1x40k)
g8516(LC-C, HC-C, hinge-CAA)Fab'-PEG(1x40k)
g8516(LC-C, HC-S, hinge-CAA)Fab'-PEG(2x20k)

Figure 6a



anti-CH1 reveal

Figure 6b



anti-Kappa reveal

Figure 7

Amino acid sequence	SEQ ID NO
Hinge sequences	
DKTHTCPP	1
DKTHTCAA	2
DKTHTCPPCPA	3
DKTHTCAACPA	4
GGGGCAA	5
DKPTCAA	6
DKPPTCAA	7
DPPPTCAA	8
SDKTHTCAA	9
SSDKTHTCAA	10
SSSDKTHTCAA	11
GCAA	12
GKTHTCAA	13
GDKTHTCAA	14
CKappa sequences	
VTKSFNRGEC	15
VTKSFNRGCS	16
VTKSFNRCES	17
VTKSFNCGES	18
VTKSFCRGES	19
VTKSCNRGES	20